SEQUENCE LISTING

<110> Acharya, et al.

<120> Crystal Structure of an Angiotensin-Converting Enzyme (ACE) and Uses Thereof

<130> 30699/41065

<160> 11

<170> PatentIn version 3.3

<210> 1
<211> 701
<212> PRT
<213> Homo sapiens

<400> 1

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Ser Gln Gln Val Thr Val Thr His Gly Thr Ser Ser Gln Ala Thr Thr
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Ser Ser Gln Thr Thr Thr His Gln Ala Thr Ala His Gln Thr Ser Ala
            20                  25                  30

Gln Ser Pro Asn Leu Val Thr Asp Glu Ala Glu Ala Ser Lys Phe Val
        35                  40                  45

Glu Glu Tyr Asp Arg Thr Ser Gln Val Val Trp Asn Glu Tyr Ala Glu
    50                  55                  60

Ala Asn Trp Asn Tyr Asn Thr Asn Ile Thr Thr Glu Thr Ser Lys Ile
65                  70                  75                  80

Leu Leu Gln Lys Asn Met Gln Ile Ala Asn His Thr Leu Lys Tyr Gly
                85                  90                  95

Thr Gln Ala Arg Lys Phe Asp Val Asn Gln Leu Gln Asn Thr Thr Ile
            100                 105                 110

Lys Arg Ile Ile Lys Lys Val Gln Asp Leu Glu Arg Ala Ala Leu Pro
        115                 120                 125

Ala Gln Glu Leu Glu Glu Tyr Asn Lys Ile Leu Leu Asp Met Glu Thr
    130                 135                 140

Thr Tyr Ser Val Ala Thr Val Cys His Pro Asn Gly Ser Cys Leu Gln
145                 150                 155                 160

Leu Glu Pro Asp Leu Thr Asn Val Met Ala Thr Ser Arg Lys Tyr Glu
                165                 170                 175
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Asp Leu Leu Trp Ala Trp Glu Gly Trp Arg Asp Lys Ala Gly Arg Ala
180 185 190

Ile Leu Gln Phe Tyr Pro Lys Tyr Val Glu Leu Ile Asn Gln Ala Ala
195 200 205

Arg Leu Asn Gly Tyr Val Asp Ala Gly Asp Ser Trp Arg Ser Met Tyr
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Glu Thr Pro Ser Leu Glu Gln Asp Leu Glu Arg Leu Phe Gln Glu Leu
225 230 235 240

Gln Pro Leu Tyr Leu Asn Leu His Ala Tyr Val Arg Arg Ala Leu His
245 250 255

Arg His Tyr Gly Ala Gln His Ile Asn Leu Glu Gly Pro Ile Pro Ala
260 265 270

His Leu Leu Gly Asn Met Trp Ala Gln Thr Trp Ser Asn Ile Tyr Asp
275 280 285

Leu Val Val Pro Phe Pro Ser Ala Pro Ser Met Asp Thr Thr Glu Ala
290 295 300

Met Leu Lys Gln Gly Trp Thr Pro Arg Arg Met Phe Lys Glu Ala Asp
305 310 315 320

Asp Phe Phe Thr Ser Leu Gly Leu Leu Pro Val Pro Pro Glu Phe Trp
325 330 335

Asn Lys Ser Met Leu Glu Lys Pro Thr Asp Gly Arg Glu Val Val Cys
340 345 350

His Ala Ser Ala Trp Asp Phe Tyr Asn Gly Lys Asp Phe Arg Ile Lys
355 360 365

Gln Cys Thr Thr Val Asn Leu Glu Asp Leu Val Val Ala His His Glu
370 375 380

Met Gly His Ile Gln Tyr Phe Met Gln Tyr Lys Asp Leu Pro Val Ala
385 390 395 400

Leu Arg Glu Gly Ala Asn Pro Gly Phe His Glu Ala Ile Gly Asp Val
405 410 415

Leu Ala Leu Ser Val Ser Thr Pro Lys His Leu His Ser Leu Asn Leu
420 425 430

Leu Ser Ser Glu Gly Gly Ser Asp Glu His Asp Ile Asn Phe Leu Met
435 440 445

Lys Met Ala Leu Asp Lys Ile Ala Phe Ile Pro Phe Ser Tyr Leu Val
450 455 460

Asp Gln Trp Arg Trp Arg Val Phe Asp Gly Ser Ile Thr Lys Glu Asn
465 470 475 480

Tyr Asn Gln Glu Trp Trp Ser Leu Arg Leu Lys Tyr Gln Gly Leu Cys
485 490 495

Pro Pro Val Pro Arg Thr Gln Gly Asp Phe Asp Pro Gly Ala Lys Phe
500 505 510

His Ile Pro Ser Ser Val Pro Tyr Ile Arg Tyr Phe Val Ser Phe Ile
515 520 525

Ile Gln Phe Gln Phe His Glu Ala Leu Cys Gln Ala Ala Gly His Thr
530 535 540

Gly Pro Leu His Lys Cys Asp Ile Tyr Gln Ser Lys Glu Ala Gly Gln
545 550 555 560

Arg Leu Ala Thr Ala Met Lys Leu Gly Phe Ser Arg Pro Trp Pro Glu
565 570 575

Ala Met Gln Leu Ile Thr Gly Gln Pro Asn Met Ser Ala Ser Ala Met
580 585 590

Leu Ser Tyr Phe Lys Pro Leu Leu Asp Trp Leu Arg Thr Glu Asn Glu
595 600 605

Leu His Gly Glu Lys Leu Gly Trp Pro Gln Tyr Asn Trp Thr Pro Asn
610 615 620

Ser Ala Arg Ser Glu Gly Pro Leu Pro Asp Ser Gly Arg Val Ser Phe
625 630 635 640

Leu Gly Leu Asp Leu Asp Ala Gln Gln Ala Arg Val Gly Gln Trp Leu
645 650 655

Leu Leu Phe Leu Gly Ile Ala Leu Leu Val Ala Thr Leu Gly Leu Ser
660 665 670

Gln Arg Leu Phe Ser Ile Arg His Arg Ser Leu His Arg His Ser His
675 680 685

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Gly Pro Gln Phe Gly Ser Glu Val Glu Leu Arg His Ser
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Leu Val Thr Asp Glu Ala Glu Ala Ser Lys Phe Val Glu Glu Tyr Asp
1               5                  10                  15

Arg Thr Ser Gln Val Val Trp Asn Glu Tyr Ala Glu Ala Asn Trp Asn
            20                  25                  30

Tyr Asn Thr Asn Ile Thr Thr Glu Thr Ser Lys Ile Leu Leu Gln Lys
        35                  40                  45

Asn Met Gln Ile Ala Asn His Thr Leu Lys Tyr Gly Thr Gln Ala Arg
    50                  55                  60

Lys Phe Asp Val Asn Gln Leu Gln Asn Thr Thr Ile Lys Arg Ile Ile
65                  70                  75                  80

Lys Lys Val Gln Asp Leu Glu Arg Ala Ala Leu Pro Ala Gln Glu Leu
                85                  90                  95

Glu Glu Tyr Asn Lys Ile Leu Leu Asp Met Glu Thr Thr Tyr Ser Val
            100                 105                 110

Ala Thr Val Cys His Pro Asn Gly Ser Cys Leu Gln Leu Glu Pro Asp
        115                 120                 125

Leu Thr Asn Val Met Ala Thr Ser Arg Lys Tyr Glu Asp Leu Leu Trp
    130                 135                 140

Ala Trp Glu Gly Trp Arg Asp Lys Ala Gly Arg Ala Ile Leu Gln Phe
145                 150                 155                 160

Tyr Pro Lys Tyr Val Glu Leu Ile Asn Gln Ala Ala Arg Leu Asn Gly
                165                 170                 175

Tyr Val Asp Ala Gly Asp Ser Trp Arg Ser Met Tyr Glu Thr Pro Ser
            180                 185                 190

Leu Glu Gln Asp Leu Glu Arg Leu Phe Gln Glu Leu Gln Pro Leu Tyr
        195                 200                 205
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Leu Asn Leu His Ala Tyr Val Arg Arg Ala Leu His Arg His Tyr Gly
    210                 215                 220

Ala Gln His Ile Asn Leu Glu Gly Pro Ile Pro Ala His Leu Leu Gly
225                 230                 235                 240

Asn Met Trp Ala Gln Thr Trp Ser Asn Ile Tyr Asp Leu Val Val Pro
                245                 250                 255

Phe Pro Ser Ala Pro Ser Met Asp Thr Thr Glu Ala Met Leu Lys Gln
            260                 265                 270

Gly Trp Thr Pro Arg Arg Met Phe Lys Glu Ala Asp Asp Phe Phe Thr
        275                 280                 285

Ser Leu Gly Leu Leu Pro Val Pro Pro Glu Phe Trp Asn Lys Ser Met
    290                 295                 300

Leu Glu Lys Pro Thr Asp Gly Arg Glu Val Val Cys His Ala Ser Ala
305                 310                 315                 320

Trp Asp Phe Tyr Asn Gly Lys Asp Phe Arg Ile Lys Gln Cys Thr Thr
                325                 330                 335

Val Asn Leu Glu Asp Leu Val Val Ala His His Glu Met Gly His Ile
            340                 345                 350

Gln Tyr Phe Met Gln Tyr Lys Asp Leu Pro Val Ala Leu Arg Glu Gly
        355                 360                 365

Ala Asn Pro Gly Phe His Glu Ala Ile Gly Asp Val Leu Ala Leu Ser
    370                 375                 380

Val Ser Thr Pro Lys His Leu His Ser Leu Asn Leu Leu Ser Ser Glu
385                 390                 395                 400

Gly Gly Ser Asp Glu His Asp Ile Asn Phe Leu Met Lys Met Ala Leu
                405                 410                 415

Asp Lys Ile Ala Phe Ile Pro Phe Ser Tyr Leu Val Asp Gln Trp Arg
            420                 425                 430

Trp Arg Val Phe Asp Gly Ser Ile Thr Lys Glu Asn Tyr Asn Gln Glu
        435                 440                 445

Trp Trp Ser Leu Arg Leu Lys Tyr Gln Gly Leu Cys Pro Pro Val Pro
    450                 455                 460
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Arg Thr Gln Gly Asp Phe Asp Pro Gly Ala Lys Phe His Ile Pro Ser
465 470 475 480

Ser Val Pro Tyr Ile Arg Tyr Phe Val Ser Phe Ile Ile Gln Phe Gln
485 490 495

Phe His Glu Ala Leu Cys Gln Ala Ala Gly His Thr Gly Pro Leu His
500 505 510

Lys Cys Asp Ile Tyr Gln Ser Lys Glu Ala Gly Gln Arg Leu Ala Thr
515 520 525

Ala Met Lys Leu Gly Phe Ser Arg Pro Trp Pro Glu Ala Met Gln Leu
530 535 540

Ile Thr Gly Gln Pro Asn Met Ser Ala Ser Ala Met Leu Ser Tyr Phe
545 550 555 560

Lys Pro Leu Leu Asp Trp Leu Arg Thr Glu Asn Glu Leu His Gly Glu
565 570 575

Lys Leu Gly Trp Pro Gln Tyr Asn Trp Thr Pro Asn Ser
580 585

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<210> 4
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